RAW SEQUENCE LISTING                    DATE: 09/15/2003
PATENT APPLICATION:  **US/09/964,858A**        TIME: 15:33:45

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Output Set: **N:\CRF4\09152003\I964858A.raw**

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 3 <110> APPLICANT: HOSTETTER, Margaret K.
 4        DEVORE-CARTER, Denise
 6 <120> TITLE OF INVENTION: ANTIBODIES TO THE PROPEPTIDE OF CANDIDA ALBICANS
 8 <130> FILE REFERENCE: P07274US02/BAS
10 <140> CURRENT APPLICATION NUMBER: US 09/964,858A
11 <141> CURRENT FILING DATE: 2001-09-28
13 <150> PRIOR APPLICATION NUMBER: US 60/237,082
14 <151> PRIOR FILING DATE: 2000-09-28
16 <160> NUMBER OF SEQ ID NOS: 8
18 <170> SOFTWARE: PatentIn version 3.1
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 1664
22 <212> TYPE: PRT
23 <213> ORGANISM: Candida albicans
25 <400> SEQUENCE: 1
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31 Leu Gln Leu Gln Pro Gln Ser Ser Ser Ala Ser Ile Phe Asn Ser Pro
32              20                  25                  30
35 Thr Lys Pro Leu Asn Phe Pro Arg Thr Asn Ser Lys Pro Ser Leu Asp
36          35                  40                  45
39 Pro Asn Ser Ser Ser Asp Thr Tyr Thr Ser Glu Gln Asp Gln Glu Lys
40      50                  55                  60
43 Gly Lys Glu Glu Lys Lys Asp Thr Ala Phe Gln Thr Ser Phe Asp Arg
44 65                  70                  75                  80
47 Asn Phe Asp Leu Asp Asn Ser Ile Asp Ile Gln Gln Thr Ile Gln His
48                  85                  90                  95
51 Gln Gln Gln Gln Pro Gln Gln Gln Gln Leu Ser Gln Thr Asp Asn
52              100                 105                 110
55 Asn Leu Ile Asp Glu Phe Ser Phe Gln Thr Pro Met Thr Ser Thr Leu
56          115                 120                 125
59 Asp Leu Thr Lys Gln Asn Pro Thr Val Asp Lys Val Asn Glu Asn His
60      130                 135                 140
63 Ala Pro Thr Tyr Ile Asn Thr Ser Pro Asn Lys Ser Ile Met Lys Lys
64 145                 150                 155                 160
67 Ala Thr Pro Lys Ala Ser Pro Lys Lys Val Ala Phe Thr Val Thr Asn
68              165                 170                 175
71 Pro Glu Ile His His Tyr Pro Asp Asn Arg Val Glu Glu Glu Asp Gln
72              180                 185                 190
75 Ser Gln Gln Lys Glu Asp Ser Val Glu Pro Pro Leu Ile Gln His Gln
76          195                 200                 205
79 Trp Lys Asp Pro Ser Gln Phe Asn Tyr Ser Asp Glu Asp Thr Asn Ala
80      210                 215                 220
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**RAW SEQUENCE LISTING**
PATENT APPLICATION: **US/09/964,858A**

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Output Set: **N:\CRF4\09152003\I964858A.raw**

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83  Ser Val Pro Pro Thr Pro Pro Leu His Thr Thr Lys Pro Thr Phe Ala
84  225                 230                 235                 240
87  Gln Leu Leu Asn Lys Asn Asn Glu Val Asn Ser Glu Pro Glu Ala Leu
88                      245                 250                 255
91  Thr Asp Met Lys Leu Lys Arg Glu Asn Phe Ser Asn Leu Ser Leu Asp
92                  260                 265                 270
95  Glu Lys Val Asn Leu Tyr Leu Ser Pro Thr Asn Asn Asn Asn Ser Lys
96              275                 280                 285
99  Asn Val Ser Asp Met Asp Ser His Leu Gln Asn Leu Gln Asp Ala Ser
100         290                 295                 300
103 Lys Asn Lys Thr Asn Glu Asn Ile His Asn Leu Ser Phe Ala Leu Lys
104 305                 310                 315                 320
107 Ala Pro Lys Asn Asp Ile Glu Asn Pro Leu Asn Ser Leu Thr Asn Ala
108                 325                 330                 335
111 Asp Ile Ser Leu Arg Ser Ser Gly Ser Ser Gln Ser Ser Leu Gln Ser
112                 340                 345                 350
115 Leu Arg Asn Asp Asn Arg Val Leu Glu Ser Val Pro Gly Ser Pro Lys
116         355                 360                 365
119 Lys Val Asn Pro Gly Leu Ser Leu Asn Asp Gly Ile Lys Gly Phe Ser
120         370                 375                 380
123 Asp Glu Val Val Glu Ser Leu Leu Pro Arg Asp Leu Ser Arg Asp Lys
124 385                 390                 395                 400
127 Leu Glu Thr Thr Lys Glu His Asp Ala Pro Glu His Asn Asn Glu Asn
128                 405                 410                 415
131 Phe Ile Asp Ala Lys Ser Thr Asn Thr Asn Lys Gly Gln Leu Leu Val
132                 420                 425                 430
135 Ser Ser Asp Asp His Leu Asp Ser Phe Asp Arg Ser Tyr Asn His Thr
136         435                 440                 445
139 Glu Gln Ser Ile Leu Asn Leu Leu Asn Ser Ala Ser Gln Ser Gln Ile
140         450                 455                 460
143 Ser Leu Asn Ala Leu Glu Lys Gln Arg Gln Thr Gln Glu Gln Glu Gln
144 465                 470                 475                 480
147 Thr Gln Ala Ala Glu Pro Glu Glu Glu Thr Ser Phe Ser Asp Asn Ile
148                 485                 490                 495
151 Lys Val Lys Gln Glu Pro Lys Ser Asn Leu Glu Phe Val Lys Val Thr
152         500                 505                 510
155 Ile Lys Lys Glu Pro Val Ser Ala Thr Glu Ile Lys Ala Pro Lys Arg
156         515                 520                 525
159 Glu Phe Ser Ser Arg Ile Leu Arg Ile Lys Asn Glu Asp Glu Ile Ala
160         530                 535                 540
163 Glu Pro Ala Asp Ile His Pro Lys Lys Glu Asn Glu Ala Asn Ser His
164 545                 550                 555                 560
167 Val Glu Asp Thr Asp Ala Leu Leu Lys Lys Ala Leu Asn Asp Asp Glu
168                 565                 570                 575
171 Glu Ser Asp Thr Thr Gln Asn Ser Thr Lys Met Ser Ile Arg Phe His
172             580                 585                 590
175 Ile Asp Ser Asp Trp Lys Leu Glu Asp Ser Asn Asp Gly Asp Arg Glu
176         595                 600                 605
179 Asp Asn Asp Asp Ile Ser Arg Phe Glu Lys Ser Asp Ile Leu Asn Asp
```

```
180     610                   615                   620
183 Val Ser Gln Thr Ser Asp Ile Ile Gly Asp Lys Tyr Gly Asn Ser Ser
184 625                   630                   635                   640
187 Ser Glu Ile Thr Thr Lys Thr Leu Ala Pro Pro Arg Ser Asp Asn Asn
188                   645                   650                   655
191 Asp Lys Glu Asn Ser Lys Ser Leu Glu Asp Pro Ala Asn Asn Glu Ser
192                   660                   665                   670
195 Leu Gln Gln Gln Leu Glu Val Pro His Thr Lys Glu Asp Asp Ser Ile
196             675                   680                   685
199 Leu Ala Asn Ser Ser Asn Ile Ala Pro Pro Glu Glu Leu Thr Leu Pro
200             690                   695                   700
203 Val Val Glu Ala Asn Asp Tyr Ser Ser Phe Asn Asp Val Thr Lys Thr
204 705                   710                   715                   720
207 Phe Asp Ala Tyr Ser Ser Phe Glu Glu Ser Leu Ser Arg Glu His Glu
208                   725                   730                   735
211 Thr Asp Ser Lys Pro Ile Asn Phe Ile Ser Ile Trp His Lys Gln Glu
212             740                   745                   750
215 Lys Gln Lys Lys His Gln Ile His Lys Val Pro Thr Lys Gln Ile Ile
216             755                   760                   765
219 Ala Ser Tyr Gln Gln Tyr Lys Asn Glu Gln Glu Ser Arg Val Thr Ser
220             770                   775                   780
223 Asp Lys Val Lys Ile Pro Asn Ala Ile Gln Phe Lys Lys Phe Lys Glu
224 785                   790                   795                   800
227 Val Asn Val Met Ser Arg Arg Val Val Ser Pro Asp Met Asp Asp Leu
228             805                   810                   815
231 Asn Val Ser Gln Phe Leu Pro Glu Leu Ser Glu Asp Ser Gly Phe Lys
232             820                   825                   830
235 Asp Leu Asn Phe Ala Asn Tyr Ser Asn Asn Thr Asn Arg Pro Arg Ser
236             835                   840                   845
239 Phe Thr Pro Leu Ser Thr Lys Asn Val Leu Ser Asn Ile Asp Asn Asp
240     850                   855                   860
243 Pro Asn Val Val Glu Pro Pro Glu Pro Lys Ser Tyr Ala Glu Ile Arg
244 865                   870                   875                   880
247 Asn Ala Arg Arg Leu Ser Ala Asn Lys Ala Ala Pro Asn Gln Ala Pro
248             885                   890                   895
251 Pro Leu Pro Pro Gln Arg Gln Pro Ser Ser Thr Arg Ser Asn Ser Asn
252             900                   905                   910
255 Lys Arg Val Ser Arg Phe Arg Val Pro Thr Phe Glu Ile Arg Arg Thr
256             915                   920                   925
259 Ser Ser Ala Leu Ala Pro Cys Asp Met Tyr Asn Asp Ile Phe Asp Asp
260     930                   935                   940
263 Phe Gly Ala Gly Ser Lys Pro Thr Ile Lys Ala Glu Gly Met Lys Thr
264 945                   950                   955                   960
267 Leu Pro Ser Met Asp Lys Asp Asp Val Lys Arg Ile Leu Asn Ala Lys
268             965                   970                   975
271 Lys Gly Val Thr Gln Asp Glu Tyr Ile Asn Ala Lys Leu Val Asp Gln
272             980                   985                   990
275 Lys Pro Lys Lys Asn Ser Ile Val Thr Asp Pro Glu Asp Arg Tyr Glu
276             995                   1000                  1005
```

```
279 Glu Leu  Gln Gln Thr Ala Ser  Ile His Asn Ala Thr  Ile Asp Ser
280      1010              1015              1020
283 Ser Ile  Tyr Gly Arg Pro Asp  Ser Ile Ser Thr Asp  Met Leu Pro
284      1025              1030              1035
287 Tyr Leu  Ser Asp Glu Leu Lys  Lys Pro Pro Thr Ala  Leu Leu Ser
288      1040              1045              1050
291 Ala Asp  Arg Leu Phe Met Glu  Gln Glu Val His Pro  Leu Arg Ser
292      1055              1060              1065
295 Asn Ser  Val Leu Val His Pro  Gly Ala Gly Ala Ala  Thr Asn Ser
296      1070              1075              1080
299 Ser Met  Leu Pro Glu Pro Asp  Phe Glu Leu Ile Asn  Ser Pro Ala
300      1085              1090              1095
303 Arg Asn  Val Ser Asn Asn Ser  Asp Asn Val Ala Ile  Ser Gly Asn
304      1100              1105              1110
307 Ala Ser  Thr Ile Ser Phe Asn  Gln Leu Asp Met Asn  Phe Asp Asp
308      1115              1120              1125
311 Gln Ala  Thr Ile Gly Gln Lys  Ile Gln Glu Gln Pro  Ala Ser Lys
312      1130              1135              1140
315 Ser Ala  Asn Thr Val Arg Gly  Asp Asp Asp Gly Leu  Ala Ser Ala
316      1145              1150              1155
319 Pro Glu  Thr Pro Arg Thr Pro  Thr Lys Lys Glu Ser  Ile Ser Ser
320      1160              1165              1170
323 Lys Pro  Ala Lys Leu Ser Ser  Ala Ser Pro Arg Lys  Ser Pro Ile
324      1175              1180              1185
327 Lys Ile  Gly Ser Pro Val Arg  Val Ile Lys Lys Asn  Gly Ser Ile
328      1190              1195              1200
331 Ala Gly  Ile Glu Pro Ile Pro  Lys Ala Thr His Lys  Pro Lys Lys
332      1205              1210              1215
335 Ser Phe  Gln Gly Asn Glu Ile  Ser Asn His Lys Val  Arg Asp Gly
336      1220              1225              1230
339 Gly Ile  Ser Pro Ser Ser Gly  Ser Glu His Gln Gln  His Asn Pro
340      1235              1240              1245
343 Ser Met  Val Ser Val Pro Ser  Gln Tyr Thr Asp Ala  Thr Ser Thr
344      1250              1255              1260
347 Val Pro  Asp Glu Asn Lys Asp  Val Gln His Lys Pro  Arg Glu Lys
348      1265              1270              1275
351 Gln Lys  Gln Lys His His His  Arg His His His His  His His Lys
352      1280              1285              1290
355 Gln Lys  Thr Asp Ile Pro Gly  Val Val Asp Asp Glu  Ile Pro Asp
356      1295              1300              1305
359 Val Gly  Leu Gln Glu Arg Gly  Lys Leu Phe Phe Arg  Val Leu Gly
360      1310              1315              1320
363 Ile Lys  Asn Ile Asn Leu Pro  Asp Ile Asn Thr His  Lys Gly Arg
364      1325              1330              1335
367 Phe Thr  Leu Thr Leu Asp Asn  Gly Val His Cys Val  Thr Thr Pro
368      1340              1345              1350
371 Glu Tyr  Asn Met Asp Asp His  Asn Val Ala Ile Gly  Lys Glu Phe
372      1355              1360              1365
375 Glu Leu  Thr Val Ala Asp Ser  Leu Glu Phe Ile Leu  Thr Leu Lys
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RAW SEQUENCE LISTING                    DATE: 09/15/2003
PATENT APPLICATION:  US/09/964,858A     TIME: 15:33:45

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Output Set: **N:\CRF4\09152003\I964858A.raw**

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379 Ala Ser  Tyr Glu Lys Pro Arg  Gly Thr Leu Val Glu Val Thr Glu
380    1385         .        1390              1395
383 Lys Lys  Val Val Lys Ser Arg  Asn Arg Leu Ser Arg Leu Phe Gly
384    1400              1405              1410
387 Ser Lys  Asp Ile Ile Thr Thr  Thr Lys Phe Val Pro Thr Glu Val
388    1415              1420              1425
391 Lys Asp  Thr Trp Ala Asn Lys  Phe Ala Pro Asp Gly Ser Phe Ala
392    1430              1435              1440
395 Arg Cys  Tyr Ile Asp Leu Gln  Gln Phe Glu Asp Gln Ile Thr Gly
396    1445              1450              1455
399 Lys Ala  Ser Gln Phe Asp Leu  Asn Cys Phe Asn Glu Trp Glu Thr
400    1460              1465              1470
403 Met Ser  Asn Gly Asn Gln Pro  Met Lys Arg Gly Lys Pro Tyr Lys
404    1475              1480              1485
407 Ile Ala  Gln Leu Glu Val Lys  Met Leu Tyr Val Pro Arg Ser Asp
408    1490              1495              1500
411 Pro Arg  Glu Ile Leu Pro Thr  Ser Ile Arg Ser Ala Tyr Glu Ser
412    1505              1510              1515
415 Ile Asn  Glu Leu Asn Asn Glu  Gln Asn Asn Tyr Phe Glu Gly Tyr
416    1520              1525              1530
419 Leu His  Gln Glu Gly Gly Asp  Cys Pro Ile Phe Lys Lys Arg Phe
420    1535              1540              1545
423 Phe Lys  Leu Met Gly Thr Ser  Leu Leu Ala His Ser Glu Ile Ser
424    1550              1555              1560
427 His Lys  Thr Arg Ala Lys Ile  Asn Leu Ser Lys Val Val Asp Leu
428    1565              1570              1575
431 Ile Tyr  Val Asp Lys Glu Asn  Ile Asp Arg Ser Asn His Arg Asn
432    1580              1585              1590
435 Phe Ser  Asp Val Leu Leu Leu  Asp His Ala Phe Lys Ile Lys Phe
436    1595              1600              1605
439 Ala Asn  Gly Glu Leu Ile Asp  Phe Cys Ala Pro Asn Lys His Glu
440    1610              1615              1620
443 Met Lys  Ile Trp Ile Gln Asn  Leu Gln Glu Ile Ile Tyr Arg Asn
444    1625              1630              1635
447 Arg Phe  Arg Arg Gln Pro Trp  Val Asn Leu Met Leu Gln Gln Gln
448    1640              1645         .    1650
451 Gln Gln  Gln Gln Gln Gln Gln  Ser Ser Gln Gln
452    1655              1660
455 <210> SEQ ID NO: 2
456 <211> LENGTH: 5194
457 <212> TYPE: DNA
458 <213> ORGANISM: Candida albicans
460 <400> SEQUENCE: 2
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463 ttaatttta ataaagaatc agtagatcta ttgttaaaag aaatgaactc aactccaagt   120
465 aaattattac cgatagataa acattctcat ttacaattac agcctcaatc gtcctcggca   180
467 tcaatattta attccccaac aaaaccattg aatttcccca gaacaaattc caagccgagt   240
469 ttagatccaa attcaagctc tgatacctac actagcgaac aagatcaaga gaaagggaaa   300
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**VERIFICATION SUMMARY**                    DATE: 09/15/2003
PATENT APPLICATION:   **US/09/964,858A**       TIME: 15:33:46

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